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(54) **Cysteine added variants of EPO and chemical modifications thereof.**

(57) Cysteine added variants ("CAVs") of EPO are provided having one or more cysteine residues substituted for selected naturally occurring amino acid residues, or inserted into the polypeptide sequence, and preferably being further modified by deletion of certain N-terminal amino acids. Such CAVs may be additionally modified by the coupling of sulfhydryl reactive compounds to the introduced cysteine residue(s) without loss of bioactivity to produce selected homogeneously modified EPO and improved pharmaceutical compositions containing the same.

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FIGURE 3a

1 10  
Met Ala Pro Pro Arg Leu Ile Cys Asp Ser Arg Val  
5' ATATG GCA CCA CCA AGA TTA ATT TGT GAT TCT AGA GTA  
TAC CGT GGT GGT TCT AAT TAA ACA CTA AGA TCT CAT

Leu Glu Arg Tyr Leu Leu Glu Ala Lys Glu Ala Glu Asn  
TTA GAA CGG TAC CTC TTG GAG GCC AAG GAG GCC GAG AAT  
AAT CTT GCC ATG GAG AAC CTC CGG TTC CTC CGG CTC TTA

25  
Ile Thr Thr Gly Cys Ala Glu His Cys Ser Leu Asn Glu  
ATC ACG ACG GGC TGT GCT GAA CAC TGC AGC TTG AAT GAG  
TAG TGC TGC CCG ACA CGA CTT GTG ACG TCG AAC TTA CTC

40  
Asn Ile Thr Val Pro Asp Thr Lys Val Asn Phe Tyr Ala  
AAT ATC ACT GTC CCA GAC ACC AAA GTT AAC TTT TAC GCG  
TTA TAG TGA CAG GGT CTC TGG TTT CAA TTG AAA ATG CGC

55  
Trp Lys Arg Met Glu Val Gly Gln Gln Ala Val Glu Val  
TGG AAA AGA ATG GAA GTT GGC CAG CAG GCT GTA GAA GTA  
ACC TTT TCT TAC CTT CAA CCG GTC GTC CGA CAT CTT CAT

70  
Trp Gln Gly Leu Ala Leu Leu Ser Glu Ala Val Leu Arg  
TGG CAG GGA TTA GCG CTA TTA AGT GAA GCT GTT CTC CGC  
ACC GTC CCT AAT CGC GAT AAT TCA CTT CGA CAA GAG GCG

85  
Gly Gln Ala Leu Leu Val Asn Ser Ser Gln Pro Trp Glu  
GGT CAG GCT TTA TTA GTC AAC TCT TCC CAG CCG TGG GAG  
CCA GTC CGA AAT AAT CAG TTG AGA AGG GTC GGC ACC CTC

100  
Pro Leu Gln Leu His Val Asp Lys Ala Val Ser Gly Leu  
CCC CTC CAG CTG CAT GTG GAT AAA GCC GTC AGT GGC CTT  
GGG GAT GTC GAC GTA CAC CTA TTT CGG CAG TCA CCG GAA

115  
Arg Ser Leu Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln  
CGC AGC CTC ACC ACT CTG CTT CGG GCT CTG GGA GCC CAG  
CGC TCG GAG TGG TGA GAC GAA GCC CGA GAC CCT CGG GTC

FIGURE 3b

Lys Glu Ala Ile Ser Pro Pro Asp Ala Ala Ser Ala Ala  
AAG GAA GCC ATC TCC CCT CCA GAT GCG GCC TCA GCT GCT  
TTC CTT CGG TAG AGG GGA GGT CTA CGC CGG AGT CGA CGA

130  
Pro Leu Arg Thr Ile Thr Ala Asp Thr Phe Arg Lys Leu  
CCA CTC CGA ACA ATC ACT GCT GAC ACT TTC CGA AAA CTC  
GGT GAG GCT TGT TAG TGA CGA CTG TGA AAG GCT TTT GAG

145  
Phe Arg Val Tyr Ser Asn Phe Leu Arg Gly Lys Leu Lys  
TTC CGA GTC TAC TCC AAT TTC CTC CGG GGA AAG CTG AAG  
AAG GCT CAG ATG AGG TTA AAG GAG GCC CCT TTC GAC TTC

160  
Leu Tyr Thr Gly Glu Ala Cys Arg Thr Gly Asp Arg  
CTG TAC ACA GGG GAG GCC TGC AGG ACA GGG GAC AGA  
GAC ATG TGT CCC CTC CGG ACG TCC TGT CCC CTG TCT

STOP  
TAA TAATGATAGGATCCT  
ATT ATTACTATCCTAGGAGATC - 5'

TECHNICAL FIELD

This invention relates generally to polypeptides modified by the attachment of compounds having  
 5 sulfhydryl reactive groups, improved methods for producing such modified polypeptides and improved  
 compositions containing them. The invention relates particularly to three modified polypeptides (IL-3, G-CSF  
 and EPO), to which sulfhydryl reactive compounds, including polymers, may be attached at selected  
 positions in the polypeptide that have been modified by the insertion of cysteine residues or the substitution  
 of cysteine residues for other residues.

10 BACKGROUND

The desirability of modifying biologically active and therapeutically useful polypeptides with a variety of  
 compounds, such as the hydrophilic polymer polyethylene glycol (PEG), to enhance their pharmacokinetic  
 properties has been noted. See, e.g., the discussion of the art in this area of polypeptide modification in  
 15 published PCT patent application WO87/00056, in U.S. Pat. No. 4,179,337, which discloses conjugating  
 water soluble polypeptides such as enzymes and insulin to PEG or PPG, and in U.S. Pat. No. 4,766,106,  
 which discloses conjugating ordinarily water insoluble beta-interferon, interleukin-2, or immunotoxins to PEG  
 homopolymers or polyoxyethylated glycerol. Such modification can reduce adverse immune response to  
 the polypeptide, increase the solubility for use in pharmaceutical preparations and maintain a desirable  
 20 circulatory level of such polypeptide for therapeutic efficacy.

One problem not addressed by the art in this area involves the extent to which a polypeptide can be  
 modified by attachment of compounds having reactive groups that will covalently bond to certain amino  
 acid residues of the polypeptide. For example, modification of a polypeptide with PEG or similar polymers,  
 can result in random attachment of the polymer at the amino terminus of the polypeptide and/or at one or  
 25 more lysine residues in the amino acid sequence of the protein. Because more than one PEG group can  
 attach to the polypeptide, the resultant composition may contain a heterogeneous mixture of "PEGylated"  
 polypeptide; some polypeptides having only one PEGylated site, others having more than one PEGylated  
 site. Such heterogeneity in composition is undesirable for pharmaceutical use. Furthermore, the non-  
 specificity with regard to the site(s) of attachment of compounds such as PEG to the polypeptide can result  
 30 in loss of biological efficacy of the polypeptide stemming from undesirable attachment to a polypeptide site  
 required for biological activity. United States Patent 4,904,584 addresses the foregoing by providing  
 materials and methods for site specific covalent modification of polypeptides by lysine insertion, removal,  
 and/or replacement. However, we have determined that the use of lysine as the attachment site for  
 modification, for example, by PEGylation, may be disadvantageous because not all modifications may result  
 35 in biologically active compounds and because steps must be taken to prevent PEGylation at N-termini in  
 cases where N-terminal PEGylation is not desired.

SUMMARY OF THE INVENTION

40 This invention provides materials and methods for site specific covalent modification of polypeptides,  
 particularly and preferably human IL-3, granulocyte colony stimulating factor (G-CSF) and erythropoietin  
 (EPO) polypeptides, permitting the production of compositions comprising homogeneously cys modified IL-  
 3s, G-CSFs and EPOs and pharmaceutical compositions containing the same. "Homogeneously cys  
 modified" as the term is used herein means substantially consistently modified only at specific, inserted or  
 45 substituted cysteine residues. A homogeneously modified IL-3 for example, includes an IL-3 composition  
 which is substantially consistently modified at position 6 (using the convention of counting from the N-  
 terminus of the mature protein) by the insertion of cysteine in place of the threonine of natural IL-3, but not  
 at other positions.

Thus, this invention first provides cysteine added variants ("CAVs") of IL-3, G-CSF and EPO. CAVs of  
 50 this invention encompass IL-3, G-CSF and EPO muteins that contain at least one additional cysteine residue  
 compared to the corresponding naturally occurring or previously known IL-3, G-CSF and EPO. The cysteine  
 residue(s) are introduced into the peptide structure of the CAVs at one or more amino acid positions in the  
 natural or previously known counterpart.

In the case of human IL-3, we have determined that the naturally occurring cysteine residues at  
 55 positions 16 and 84 form a disulfide bridge, essential to preserving the desired biological activity of the  
 polypeptide. For the addition of novel cysteines, some positions within the polypeptide, such as position 15  
 and 51 are unsuitable; cysteines introduced at these positions give rise to human IL-3 polypeptides with  
 substantially reduced biological activity. However, certain substitutions or deletions of residues 1-14 do not

significantly diminish the desired biological activity of IL-3. Therefore, a preferred region of novel cysteine introduction into the polypeptide is within positions 1-14 inclusive. Currently, positions 6-12 inclusive are especially preferred sites for cysteine introduction. The subsequent attachment of sulfhydryl reactive compounds, including polymers, as discussed below, to the novel cysteines added at selected positions within this region will not result in any significant loss of biological activity.

By "cysteine added variant" as the term is used herein, we mean variants of IL-3, G-CSF and EPO that are modified in amino acid structure relative to naturally occurring or previously known counterparts such that at least one cysteine residue is inserted into the natural or previously known sequence and/or is used to replace a different amino acid within that sequence.

Additionally, with respect to IL-3, the native or "natural" IL-3 sequence, with an added initiator methionine for bacterial expression, may be further modified such that the first alanine is deleted at the N-terminus of the mature polypeptide, altering the amino terminal sequence from MET\*ALA\*PRO to MET\*PRO (the "mp" mutein). For the "mp" mutein, such N-terminus modification permits more consistent removal of the N-terminal methionine. As is already known, in bacterial expression systems, cleavage at the N-terminal methionine occurs. Likewise, the native EPO N-terminal sequence (with the added MET) begins MET\*ALA\*PRO and it may prove advantageous to delete the first alanine to obtain an mpEPO mutein. With regard to G-CSF, the natural human N-terminal sequence begins with MET\*THR\*PRO (with the MET added for bacterial production) and it may be desirable to delete this N-terminal threonine to advantageously obtain a mpG-CSF mutein.

Alternatively, the natural IL-3 sequence may be further modified such that the first two amino acids at the N-terminus of the mature polypeptide are deleted, leaving a terminus beginning with MET\*THR\*GLN\*THR (the "m3" mutein). For the "m3" mutein, such N-terminus modification permits one to take advantage of the methionine at position 3 in the naturally occurring human IL-3 molecule, as the initiator methionine.

The CAVs of this invention make it possible to produce homogeneous, biologically active IL-3, G-CSF and EPO compositions substantially specifically and consistently modified at selected positions with sulfhydryl reactive compounds (described hereinafter).

In the practice of this invention, at least one cysteine residue is introduced in that portion of the IL-3, G-CSF or EPO polypeptide where modification via a sulfhydryl reactive compound is desired. The cysteine residue or residues are so introduced by genetic engineering methods as described below. Novel cysteine residues may be engineered into the polypeptide for example, by simple insertion of a cysteine codon into the DNA molecule at the desired site or by converting a desirably located asparagine or other codon to a cysteine codon. Convenient methods for site specific mutagenesis or DNA synthesis for producing a DNA molecule encoding the desired CAV, expression in procaryotic or eucaryotic host cells of the DNA molecule so produced, and recovery of the CAV produced by such expression are also disclosed.

The CAVs of this invention retain useful biological properties of the natural or previously known protein and may thus be used for applications identified for the non-modified parent. Modification with such sulfhydryl reactive compounds, however, is preferred. Such biologically active, modified CAVs can be produced in homogeneous compositions which, it is contemplated, will provide improved pharmacokinetic profiles, immunogenicity profiles, and/or solubility characteristics relative to the parent polypeptides. Furthermore, CAVs may enable the formation of multimeric forms of the normally monomeric polypeptide with the same, albeit improved characteristics. Multimeric CAVs also enable the formation of "hetero-conjugates"—i.e., two or more distinct polypeptides joined via the sulfhydryl groups of the added cysteine residues, e.g., IL-3 joined to EPO or IL-3 joined to G-CSF.

Biological activity of the CAVs before or after modification with the sulfhydryl reactive compounds may be determined by standard *in vitro* or *in vivo* assays conventional for measuring activity of the parent polypeptide. Alternatively, we provide herein a "small scale" screening method wherein successful Cys modification and attachment of the sulfhydryl reactive compound may be tested.

Selective and homogeneous modification of the CAVs with sulfhydryl reactive compounds is possible since such compounds will covalently bond primarily only to the cysteine residue(s) in the CAV. Secondary reactivity at His, Lys and Tyr residue(s) may be observed, depending on the choice of sulfhydryl reactive compound, but at a significantly lower rate. The modified CAVs so produced may then be recovered, and if desired, further purified and formulated into pharmaceutical compositions by conventional methods.

Sulfhydryl reactive compounds include compounds such as polyalkylene glycol, e.g. polyethylene and polypropylene glycol, as well as derivatives thereof, with or without coupling agents or derivatization with coupling or activating moieties, for example, with thiol, triflate, tresylate, aziridine or oxirane, or preferably with S-pyridyl or maleimide moieties. Compounds such as S-Pyridyl Monomethoxy PEG and Maleimido Monomethoxy PEG are exemplary. Additionally, sulfhydryl reactive compounds include, but are not limited

to, charged or neutral polymers of the following types: dextran, colominic acids or other carbohydrate based polymers, polymers of amino acids and biotin derivatives, resulting in a protein modified with this well known affinity reagent often used for antibody based assays.

Briefly, the method comprises reacting the CAV with a sulfhydryl reactive compound under suitable conditions, preferably non-denaturing conditions, and in sufficient amounts permitting the covalent attachment of the sulfhydryl reactive compound to the introduced cysteine residue(s) present in the polypeptide backbone of the CAV. The reaction may be reducible or non-reducible; and generally, the amount of sulfhydryl reactive compound used should be at least equimolar to the number of cysteines to be derivatized, although use of excess sulfhydryl reactive compound is preferred, both to improve the rate of reaction and to insure consistent modification at all reactive sites. The modified CAV produced, may then be recovered, purified and formulated by conventional methods. See e.g., WO 87/00056 and references cited therein.

Other aspects of the present invention include therapeutic methods of treatment and therapeutic compositions which employ the modified CAVs of the present invention, either alone or with other lymphokines, hematopoietins and/or growth factors, such as granulocyte macrophage colony-stimulating factor (GM-CSF), macrophage colony-stimulating factor (M-CSF), IL-1, IL-2, IL-4, IL-5, IL-6 and IL-10. These methods and compositions take advantage of the improved pharmacokinetic properties of these modified CAVs to provide treatments, e.g., such as employing lower dosages of polypeptide, less frequent administration, lower immunogenicity and more desirable distribution, required for the therapeutic indications for the natural polypeptide.

Other aspects and advantages of the present invention will be apparent upon consideration of the following detailed description of the invention, including illustrative examples of the practice thereof.

#### BRIEF DESCRIPTION OF THE DRAWINGS

Fig. 1 is the human IL-3 gene construct for *E. coli* expression, having the polypeptide sequence shown of natural (wild type) human IL-3, plus an initiator methionine, as expressed in *E. coli*, with the amino acids numbered from the N-terminus for reference to the muteins discussed herein.

Fig. 2 is the human G-CSF gene construct for *E. coli* expression, having the polypeptide sequence shown of natural (wild type) human G-CSF, plus an initiator methionine, as expressed in *E. coli* with the amino acids numbered from the N-terminus for reference to the muteins discussed herein.

Fig. 3 is a chemically synthesized human EPO gene construct for *E. coli* expression, having the polypeptide sequence of natural (wild type) human EPO, plus an initiator methionine, as expressed in *E. coli* with the amino acids numbered from N-terminus for reference to the muteins discussed herein.

#### DETAILED DESCRIPTION OF THE INVENTION

The present invention involves the selective modification of IL-3, G-CSF and EPO for pharmaceutical use, to both enhance their pharmacokinetic properties and provide homogeneous compositions for human therapeutic use. Although human IL-3, DNA and peptide sequences are preferred as the starting point in this invention as it relates to IL-3, any primate IL-3 is susceptible to use in the method of the invention, given the significant homology between e.g., human and gibbon species of the protein and DNA. See Leary et al., Blood (1982) 70: 1343-1348. The method for selectively modifying IL-3, G-CSF and EPO involves selecting locations in the polypeptide sequence for the attachment of sulfhydryl reactive compounds. This step may be accomplished by altering the amino acid sequence of the polypeptide by inserting cysteine residues at selected sites or by converting selected endogenous residues into cysteine residues. For example, the codons AAA or AAG, which code for lysine, can be changed to the codon TGC or TGT, which code for cysteine.

CAVs in accordance with this invention also include allelic variations in the protein sequence, i.e. sequence variations due to natural variability from individual to individual, or with other amino acid substitutions or deletions which still retain desirable biological properties of the parent.

All CAVs of this invention may be prepared by expressing recombinant DNA sequences encoding the desired variant in host cells, e.g. procaryotic host cells such as *E. coli*, or eucaryotic host cells such as yeast or mammalian host cells, using methods and materials, e.g. vectors, as are known in the art. Host cells containing and capable of expressing the CAV-encoding DNA are thus encompassed by this invention. DNA sequences encoding the variants may be produced synthetically or by conventional site-directed mutagenesis of DNA sequences encoding the protein or polypeptide or analogs thereof. Figure 1 shows the human IL-3 gene construct inserted in plasmid pAL-hIL3-781 and expressed in the *E. coli* K12 strain

designated GI586. This strain containing the plasmid was deposited with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland 20852 USA on April 19, 1989 and given accession number 67932. Other DNA sequences for natural primate IL-3 have been cloned and the DNA sequences, including cDNA sequences, and specific peptide sequences for the same have been published, in PCT application number US87/01702, published as WO 88/00598 on January 28, 1988, and are therefore known in the art. These DNA sequences have been deposited with the American Type Culture Collection and given accession numbers ATCC 67154, 67326, 67319 and its replacement 68042, and 40246. DNA sequences for natural G-CSF and EPO have been cloned and the sequences and their corresponding peptide sequences published and are therefore known in the art.

DNA molecules encoding natural human IL-3s, G-CSFs and EPOs therefore may be obtained (i) by cloning in accordance with the published methods, (ii) from the deposited plasmids, or (iii) by synthesis, e.g. using overlapping synthetic oligonucleotides based on the published sequences which together span the desired coding region. Such methods are known in the art. See the foregoing PCT application published as WO 88/00598 and PCT application number US88/00402 published as WO88/06161.

As mentioned above, DNA sequences encoding individual CAVs of this invention may be produced synthetically or by conventional site-directed mutagenesis of a DNA sequence encoding the parental polypeptides or analogs thereof. Such methods of mutagenesis include the M13 system of Zoller and Smith, *Nucleic Acids Res.* (1982) 10:6487 - 6500; *Methods Enzymol.* (1983) 100:468-500; and *DNA* (1984) 3:479-488, which uses single stranded DNA and the method of Morinaga et al., *Bio/technology* (July 1984) 636-639, which uses heteroduplexed DNA. Exemplary oligonucleotides used in accordance with such methods are described below. It should be understood, of course, that DNA encoding each of the CAVs of this invention may be analogously produced by one skilled in the art through site-directed mutagenesis using appropriately chosen oligonucleotides.

The new DNA sequences encoding the CAVs of this invention can be introduced into appropriate vectors for heterologous expression in the desired host cells, whether procaryotic or eucaryotic. The activity produced by the transiently transfected or stably transformed host cells (or their progeny) may be measured by using standard assays conventional for the parental protein. Where the host cell is bacterial, the DNA should be free of introns, e.g. a cDNA or synthetic DNA, and may be free of any secretory leader sequence. For eucaryotic expression, introns may be present or absent and a secretory leader sequence should preferably be present.

The CAVs produced by expression in the genetically engineered host cells may then be purified, and if desired formulated into pharmaceutical compositions by conventional methods, often preferably by methods which are typically used in purifying and/or formulating the parental protein. It is contemplated that such pharmaceutical compositions containing the CAV in admixture with a pharmaceutically acceptable carrier will possess similar utilities to those of the parental proteins, such as those set forth in WO 88/00598 *supra*, at page 3.

In another, and preferred, aspect of this invention, the CAVs produced by recombinant means as mentioned above are reacted with the desired sulfhydryl reactive compound under conditions permitting attachment of the sulfhydryl reactive moiety to the sulfhydryl group of the introduced cysteine residues in the peptide backbone of the CAV. These modified CAVs, preferably produced initially on a small scale, may then be screened for bioactive muteins possessing the sulfhydryl reactive compounds attached to the site or sites desired. Alternatively, this screening may be accomplished before attachment with the sulfhydryl reactive compound.

The term "sulfhydryl reactive compound" is defined herein as any compound having, or capable of being activated to have, a reactive group capable of forming a covalent attachment to the sulfhydryl group (-SH) of the cysteine residue. Included among such compounds are polymers such as PEG and polypropylene glycol (PPG), dextran, colominic acids or other carbohydrate based polymers and polymers of amino acids and biotin derivatives. Activation may occur by modification of the compound with a sulfhydryl moiety, such as a sulfhydryl group, thiol, triflate, tresylate, aziridine or oxirane, or preferably, with S-pyridyl or maleimide. The sulfhydryl reactive compound need not have any particular molecular weight, but a molecular weight of between about 1,000 and 30,000 for the activated compound is preferred, especially for PEG. Methods of attachment will be described in detail below. By controlling the number and location of the cysteines in the CAV sequence, the number and location(s) of the attached sulfhydryl reactive compound can be selectively controlled. Such control of attachment location and number enables the production of only certain selectively modified molecules retaining the desired biological activity, rather than production of a heterogeneous mixture of variably modified molecules, only some of which may be active. It is also important to note that this positional selectivity of the PEGylation or other attachment allows the normal functional interactions of the protein to be preserved, blocked, or regenerated by release of the sulfhydryl

reactive compound.

Another aspect of the invention is therefore homogeneous compositions of modified CAVs as described herein, e.g. PEGylated CAVs. Specific embodiments of IL-3 CAVs of the invention include human IL-3 which has a cysteine residue replacing lysine at position 10 and the m3 initiation sequence. (Amino acid numbers  
5 for the CAVs of the present invention are used herein in the conventional manner, sequentially from the N-terminus, and correlate with the numbering system used in Fig. 1 for the natural human IL-3 as expressed in *E. coli*.) Similarly, the naturally occurring lysine residue in human IL-3 (Fig. 1) at amino acid position 100 may be converted to a cysteine to create a human IL-3 CAV of the invention. Another embodiment has cysteine at positions 9 and 10 and the m3 initiation sequence.

10 Specific embodiments of G-CSF and EPO CAVs of the invention include human G-CSF which has an alanine residue replacing the naturally occurring cysteine residue at position 17 and a cysteine residue replacing the naturally occurring alanine at position 37, and human EPO which has a cysteine residue replacing serine at position 9. The modification of Ala17 of G-CSF is made to prevent possible improper disulfide bridge formation.

15 For bacterial expression where the secretory leader-encoding DNA sequence is removed from the CAV-encoding DNA, it may be desirable to additionally modify the sequence such that it encodes an N-terminus comprising Met-Pro— (the mp mutein) instead of other N-termini such as Met-Ala-Pro (in IL-3 and EPO) or Met-Thr-Pro (in G-CSF). Such N-terminal modification permits more consistent removal of the N-terminal methionine. Alternatively, the first two residues of natural, human IL-3 may be deleted, leaving the naturally  
20 occurring methionine at position 3 as the translation initiator (the m3 mutein).

CAVs of this invention, modified as described, encompass CAVs containing other modifications as well, including truncation of the peptide sequence, deletion or replacement of additional amino acids with amino acids other than cysteine, insertion of new N-linked glycosylation sites, abolishment of natural N-linked glycosylation sites, etc., so long as the bioactivity of the molecule is retained. Thus, this invention  
25 encompasses CAVs encoded for by DNA molecules which are capable of hybridizing under stringent conditions to the DNA molecule encoding the parental IL-3, G-CSF or EPO (or would be so capable but for the use of synonymous codons) so long as the encoded polypeptide contains one or more additional introduced cysteine residues relative to the parental peptide sequence. Exemplary stringent conditions can be found in T. Maniatis, et al., *Molecular Cloning (A Laboratory Manual)*, Cold Spring Harbor Laboratory  
30 (1982), pages 387-389. An example of one such stringent hybridisation condition is 4XSSC at 65°C, followed by a washing in 0.1XSSC at 65°C for an hour. Alternatively, an exemplary stringent hybridization condition is 50% formamide, 4XSSC at 42°C.

Because the method and compositions of this invention provide homogeneous modified IL-3s, G-CSFs and EPOs, the invention also encompasses such homogeneous compositions for pharmaceutical use which  
35 comprise a therapeutically effective amount of a modified CAV described above in admixture with a pharmaceutically acceptable carrier. Such composition can be used in generally the same manner as that described for the natural or recombinant polypeptides. It is contemplated that the compositions will be used for treating a variety of conditions, e.g. involving stimulating hematopoiesis or improving a patient's hematological profile. For example, a modified IL-3 of the present invention may be used as an adjunct to  
40 cancer chemotherapy, radiation therapy, or in the treatment of immune disorders, as discussed in WO 88/00598, at page 17-19. The exact dosage and method of administration will be determined by the attending physician depending on the particular modified CAV employed, the potency and pharmacokinetic profile of the particular compound as well as on various factors which modify the actions of drugs, for example, body weight, sex, diet, time of administration, drug combination, reaction sensitivities and severity  
45 of the particular case. Generally, the daily regimen should be in the range of the dosage for the natural or recombinant unmodified protein, e.g. a range of about 0.1 to about 100 µg of polypeptide per kilogram of body weight, preferably from about 0.1 to about 30 µg of polypeptide per kilogram of body weight.

The therapeutic method and compositions of the present invention may also include co-administration with other drugs or human factors. A non-exclusive list of other appropriate hematopoietins, CSFs (colony  
50 stimulating factors) and interleukins for simultaneous or serial co-administration with the CAVs of the present invention includes GM-CSF, CSF-1 (in its various known forms; CSF-1 is also referred to as M-CSF or macrophage colony-stimulating factor), Meg-CSF, IL-1, IL-2, IL-4, IL-6, IL-10, B-cell growth factor, B-cell differentiation factor and eosinophil differentiation factor. Additionally, the CAVs of the present invention may be administered with, or chemically attached to, monoclonal or polyclonal antibodies in a therapeutic use.  
55 Alternatively, these growth factors may be attached to certain toxins, e.g., ricin, for use in a therapeutic regimen. The dosage recited above would be adjusted to compensate for such additional components in the therapeutic composition or regimen. In the case of pharmaceutical compositions containing modified lymphokine CAVs, for example, progress of the treated patient can be monitored by periodic assessment of

the hematological profile, e.g. white cell count, hematocrit and the like.

The following examples illustratively describe the CAVs and the methods and compositions of the present invention.

## 5 EXPERIMENTAL MATERIALS, METHODS AND EXAMPLES

### EXAMPLE 1: Eucaryotic Expression Materials and Methods

Eukaryotic cell expression vectors into which DNA sequences encoding CAVs of this invention may be  
 10 inserted (with or without synthetic linkers, as required or desired) may be synthesized by techniques well known to those skilled in this art. The components of the vectors such as the bacterial replicons, selection genes, enhancers, promoters, and the like may be obtained from natural sources or synthesized by known procedures. See Kaufman et al., J. Mol. Biol., (1982) 159: 601-621, Kaufman, Proc. Natl. Acad. Sci. (1985) 82:689-693. See also WO 87/04187, filed January 2, 1987 (pMT2 and pMT2-ADA), and US Patent  
 15 Application Serial No. 88,188, filed August 21, 1987 (pxMT2). Exemplary vectors useful for mammalian expression are also disclosed in the patent applications cited in Example 4, which are hereby incorporated by reference. Eucaryotic expression vectors useful in producing variants of this invention may also contain inducible promoters or comprise inducible expression systems as are known in the art. See US Patent Application Serial No. 893,115 (filed August 1, 1986) and PCT/US87/01871, published as WO88/00975 on  
 20 February 11, 1988.

Established cell lines, including transformed cell lines, are suitable as hosts. Normal diploid cells, cell strains derived from *in vitro* culture of primary tissue, as well as primary explants (including relatively undifferentiated cells such as hematopoietic stem cells) are also suitable. Candidate cells need not be genotypically deficient in the selection gene so long as the selection gene is dominantly acting.

25 If eucaryotic host cells are used, they will preferably will established mammalian cell lines. For stable integration of the vector DNA into chromosomal DNA, and for subsequent amplification of the integrated vector DNA, both by conventional methods, CHO (Chinese Hamster Ovary) cells are presently preferred in such embodiments. Alternatively, the vector DNA may include all or part of the bovine papilloma virus genome (Lusky et al., Cell (1984) 36: 391-401) and be carried in cell lines such as C127 mouse cells as a  
 30 stable episomal element. Other usable mammalian cell lines include HeLa, COS-1 monkey cells, melanoma cell lines such as Bowes cells, mouse L-929 cells, 3T3 lines derived from Swiss, Balb-c or NIH mice, BHK or HaK hamster cell lines and the like.

Stable transformants then are screened for expression of the CAV product by standard immunological or activity assays. The presence of the DNA encoding the CAV polypeptides may be detected by standard  
 35 procedures such as Southern blotting. Transient expression of the CAV genes during the several days after introduction of the expression vector DNA into suitable host cells such as COS-1 monkey cells is measured without selection by activity or immunologic assay of the proteins in the culture medium.

Following the expression of the DNA by conventional means, the CAVs so produced may be recovered, purified, and/or characterized with respect to physicochemical, biochemical and/or clinical parameters, all by  
 40 known methods.

### EXAMPLE 2: Bacterial and Yeast expression

Bacterial and yeast expression may be effected by inserting (with or without synthetic linkers, as  
 45 required or desired) the DNA molecule encoding the desired CAV into a suitable vector (or inserting the parental DNA sequence into the vector and mutagenizing the sequence as desired therein), then transforming the host cells with the vector so produced using conventional vectors and methods as are known in the art, e.g. as disclosed in published PCT Application No. WO 86/00639, published January 30, 1986. Transformants are identified by conventional methods and may be subcloned if desired. Characterization of  
 50 transformants and recombinant product so produced may be effected and the product recovered and purified, all as described in Example 1.

For bacterial expression, the DNA sequences encoding the CAVs are preferably modified by conventional procedures to encode only the mature polypeptide and may optionally be modified to include preferred bacterial codons.

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Expression in E. coli:

The IL-3 CAVs of Example 5 were expressed in *E. coli* as follows: Plasmid pAL-hIL3-781 was transformed into an *E. coli* K12 strain GI586, a derivative of strain W3110 in which the C<sub>I</sub> and Rex regions of bacteriophage lambda carrying the C857 allele have been inserted into the ClaI site of the lacZ gene of the bacterial genome. This insert consists of all of the DNA sequences between nucleotides 35711 and 38104 of the phage genome. See F. Sanger, et al., *J. Mol. Biol.* (1982) 162:729. *E. coli* K12 strain GI586 (pAL-hIL3-781) was deposited at the ATCC on April 19, 1989 and given accession number 67932.

When GI586 transformed with pAL-hIL3-781 is grown at 30 degrees centigrade to high cell density and then heated to 40 degrees centigrade, IL-3 is produced rapidly and accumulates over the next two or three hours to reach greater than 10 percent of the total cellular protein. This protein is produced in an insoluble form which must be solubilized and refolded by conventional methods. See, e.g., T.E. Creighton, *Prog. Biophys. Molec. Biol.* (1978) 33:231-297. Following expression, the CAVs so produced were recovered, purified and characterized as follows.

The G-CSF CAV of Example 5 and the EPO CAV of Example 6 were expressed similarly in *E. coli* by removing the DNA sequence encoding CAV IL-3 from pAL-hIL3-781 and inserting the appropriate G-CSF or EPO DNA sequence as set forth in those Examples. Transformations were carried out under the same conditions set forth above.

1. Purification of CAV IL-3

All buffers were prepared using glass distilled water; all were degassed for at least five minutes, using house vacuum/sonification, prior to the addition of DTT.

First, 400 grams wet weight frozen *E. coli* cell paste was suspended in 2500 ml of buffer containing 50 mM Tris-HCl, pH 8.5, 1 mM EDTA, 5 mM P-aminobenzamidine, 1 mM PMSF and 2 mM DTT (hereinafter in this Example "buffer A"), to obtain a final volume of 2850 ml. Glass rods and magnetic stirrers were used to resuspend the cell paste. Then the cell suspension was lysed by passing it through a matin gaulin valve at 9000 psi four times, with cooling between each time. Temperature was maintained below 30 degrees centigrade by collection of the lysate into glass vessels cooled in ice/water mixture. Protein concentration was 22 mg/ml; final volume was 2850 ml.

The lysate was centrifuged for 30 minutes at 8000 rpm in a Sorval centrifuge with a GS-3 rotor. The supernatant (2600 ml at 17.0 mg/ml) was discarded and the resultant pellet (hereinafter in this Example P1) from this centrifugation was resuspended in approximately 400 ml buffer A, using glass rods and a magnetic stirrer. The milky suspension was then passed through an 18 gauge needle using a 60 ml syringe. The final volume was 640 ml, with a protein concentration of 25.6 mg/ml.

The resuspended P1 pellet was then centrifuged for 10 minutes in a Sorval centrifuge with a GS-3 rotor at 8000 rpm. The supernatant from this centrifugation was poured into two fresh centrifuge tubes (hereinafter in this example "S2") and the resultant pellet ("P2") was resuspended in buffer A to a final volume of 165 ml, with a protein concentration of 50 mg/ml. The S2 supernatant was then centrifuged for 10 minutes and the resulting pellet ("P3") was resuspended in 65 ml buffer A. The resulting supernatant ("S3") was further centrifuged for 10 minutes and the resulting P4 pellet was resuspended in buffer A to a final volume of 50 ml, with a protein concentration of 11.3 mg/ml. Because the P4 pellet contained so little IL-3, it was not used in subsequent steps. The S4 supernatant from the final centrifugation, approximately 600 ml, contained the membranous components at a concentration of approximately 10 mg/ml.

The P2 and P3 pellets were pooled and centrifuged at 9000 rpm (GSA rotor) for 10 minutes yielding two pellets ("P2-2") and a cloudy supernatant, which using HPLC analysis was found void of IL-3 and was discarded. The P2-2 pellet was frozen at -20 degrees centigrade for later use.

The frozen P2-2 pellet was then resuspended in buffer A (which contained 10 mM DTT rather than 2 mM DTT) to a final volume of 100 ml using glass rods and magnetic stirrer and then passed through an 18 gauge needle. 400 ml of 7 M fresh guanidine in the 10 mM DTT buffer A was added to the resuspended P2-2 and after one quick inversion, the solubilized P2-2 pellet was immediately placed in 3 x 250 centrifuge tubes and centrifuged for 15 minutes at 8000 rpm (GSA rotor). 500 ml of the supernatant at a concentration of 5.98 mg/ml was purified further at room temperature by RP-HPLC. The foregoing two steps were performed in 17 to 22 minutes.

This purification protocol may be applied to the purification of the G-CSF and EPO CAVs expressed in *E. coli* as set forth above with similar results.

## 2. RP-HPLC separation of IL-3 CAVs

The buffers used in this separation protocol were 0.1% (v/v) TFA in water, and 0.1% TFA in acetonitrile.

A two inch Vydac C4 column was equilibrated in 10% acetonitrile. The supernatant from the 7 M guanidine solubilization was immediately applied onto the C4 column having a volume of approximately 470 ml at 180 ml per minute. The column was developed at 20 ml per minute and was washed in 10% acetonitrile until absorbance at 280 nm was back to baseline. The following gradient was established by washing with the following concentrations of acetonitrile at the following times:

time (in minutes)	% acetonitrile
5	10
10	35
55	55
60	80
65	80
67.5	10

40 ml fractions were collected after 35 minutes into the gradient. 10  $\mu$ l samples were removed from each fraction, vacuum speed dried and taken up in 20  $\mu$ l of 2x SDS-sample Laemmli buffer. SDS-PAGE analysis was performed and IL-3 presence was confirmed. All fractions were then frozen at -80 degrees centigrade

Similarly, RP-HPLC separation of G-CSF and EPO CAVs may be accomplished.

## 3. Refolding of IL-3 CAVs

One of the RP-HPLC separated fractions containing approximately 75 mg (7.5 ml) IL-3 was diluted to approximately 0.5 mg/ml by the addition of 142.5 ml of 6.4 M guanidine in 50 mM NaPO<sub>4</sub> pH 7.0, 1 mM EDTA and 0.2 mM DTT. The mixture was then added to 750 ml of 50 mM NaPO<sub>4</sub> pH7, 1 mM EDTA, 0.2 mM DTT buffer, transferred to dialysis tubing and dialyzed for two hours against 4 L of the same buffer. The IL-3 (now approximately 0.22 M guanidine) was twice further dialyzed against 8 L of the same buffer containing 0.1 mM DTT.

PEGylation of this purified IL-3 is set forth in Examples 7 and 8 below.

Refolding of G-CSF and EPO CAVs may be accomplished in the same manner.

## 4. Confirmation of bioactivity

Bioactivity of IL-3 CAVs or EPO CAVs may be confirmed by using the TF-1 cell proliferation assay. The TF-1 cell line has been described (Kitamura et al., Blood (1989) 73:375-380).

In that assay, cells are maintained at 37 degrees centigrade in humid air containing 5% CO<sub>2</sub> and culture media used is RPMI (Gibco), 10% heat inactivated fetal calf serum, 2 mM L-glutamine, with 5 ng/ml of recombinant GM-CSF added. Every 3-4 days cells are adjusted to a density of 2 x 10<sup>5</sup> cells/ml. Just prior to assay the cells are centrifuged 500xG, 5 minutes, washed in culture media without rGM-CSF, recentrifuged and resuspended at a density of 10<sup>5</sup> cells/ml.

IL-3 or EPO samples to be assayed are diluted between 1:500 and 1:10,000 in culture media without rGM-CSF. 125  $\mu$ l of the diluted sample is placed in the top row of a 96 well microtiter plate. The remaining wells are filled with 100  $\mu$ l of culture media without rGM-CSF and the top row samples are serially diluted five fold down the microtiter plate. To each well, 100  $\mu$ l of diluted cells (10<sup>4</sup> cells) are added and the plate is incubated at 37 degrees centigrade, 5% CO<sub>2</sub> for 48-72 hours. Thereafter, 0.5 uCi <sup>3</sup>H-thymidine is added per well and the plate is further incubated for 4-6 hours. Cells are then harvested using an automated cell harvester (LKB 1295-001) and the <sup>3</sup>H-thymidine uptake is quantitated.

Alternatively for IL-3, a CML proliferation assay as described in PCT/US87/017024, International Publication Number WO88/00598, published January 28, 1988, can be used.

Bioactivity of G-CSF CAVs may be confirmed using the 32D murine cell line, as described in Hapel, et al., Blood (1984) 64:786-790, and adding 5% v/v of WEHI-3 conditioned media from a 48 hour culture of WEHI-3 cells, ATCC TIB68, (1x10<sup>6</sup> cells/ml) in RPMI-1640 media supplemented with 2 mM L-glutamine instead of the 5 ng/ml recombinant GM-CSF in the TF-1 cell proliferation protocol. The wash and assay steps are then carried out in the absence of WEHI-3 conditioned media.

EXAMPLE 4: Mutagenesis Protocol

Site directed mutagenesis may be effected using conventional procedures known in the art. See e.g., International Applications Nos. WO 87/07144, and WO 87/04722, and US Patent Application Serial Nos.  
5 099,938 (filed September 23, 1987) and 088,188 (filed August 21, 1987) and the references cited therein.

EXAMPLE 5: Exemplary Mutagenesis Reactions

The following human IL-3, G-CSF and EPO muteins were engineered by substitution of the codons  
10 indicated for a cys codon, or by insertion of a cys codon, using conventional site directed mutagenesis techniques:

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IL-3

	<u>mp mutein</u>	<u>m3 mutein</u>	<u>cys modification</u>
5	mpCys10	m3cys10	AAA to TGC (Lys to Cys)
	mpCys6	m3Cys6	ACT to TGC (Thr to Cys)
10	mpCys8	m3Cys8	TCT to TGC (Ser to Cys)
	mpCys12	m3Cys12	TCT to TGC (Ser to Cys)
	mpCys100	m3Cys100	AAG to TGT (Lys to Cys)
15	mpCys134	m3Cys134	Insertion of TGT between TTC and TAG (Cys between Phe 133 and stop codon)
	mpCys3		ATG to TGC (Met to Cys)
20	mpAlCys19		Replacement of amino acids 1-15 with the "mp" terminus and modif. of pos. 19 from ATG to TGC (Met to Cys)
25		m3Cys6,10	ACT and AAA to TGC (Thr and Lys to Cys)
30		m3Cys9,10	TTA and AAA to TGC (Leu and Lys to Cys)
		m3Cys6,8	ACT and TCT to TGC (Thr and Ser to Cys)
35		m3Cys6,8,10	ACT, TCT and AAA to TGC (Thr, Ser and Lys to Cys)
40		m3Cys8,9,10	TCT, TTA and AAA to TGC (Ser, Leu and Lys to Cys)

G-CSF

	<u>mp mutein</u>	<u>cys modification</u>
45	mpAla17Cys37	GCC to TGC (Ala to Cys)

EPO

	<u>mp mutein</u>	<u>cys modification</u>
50	mpCys9	TCT to TGT (Ser to Cys)

In the examples depicted above the modification site of the natural IL-3, G-CSF or EPO protein is designated by the number after "Cys" and the amino acid sequence of the CAV is identical to that of the native protein, except for the position indicated, with respect to the N-terminus (see Fig. 1). The "mp" and "m3" designations signify the two different alterations of the N-terminus that will be discussed in detail below. Additionally, cys may be introduced in place of native IL-3 codons, for example at positions 63 or 66,

alone or in combination with other cys introduction(s), e.g. at position 10--with any of the described N-termini. Contemplated EPO muteins include EPO mpCys166, having the mp N-terminus and the native arginine at position 166 deleted and replaced with cysteine, and mpCys24Cys38Cys83, having the mp N-terminus and the three N-linked glycosylation sites at the asparagine amino acids replaced with cysteines.

5 With respect to IL-3 muteins, certain point modifications may result in partial loss of biological activity or inability of the sulfhydryl reactive compound to attach. For example, modification at position 28 results in a biologically active CAV, but attachment of a sulfhydryl reactive compound fails, possibly because position 28 appears internally in the refolded CAVs tertiary structure. Compare, Wingfield, D., et al., Eur. J. Biochem. (1989) 179:565-571, in which the authors discussed the Cys modification of IL-1 $\beta$  at position 138 to active  
10 IL-1 $\beta$ -phycoerythrin conjugate. Additionally, we have found that substitution of a cysteine residue for the amino acids at positions 15 or 51 of the natural human IL-3 may result in partial loss of bioactivity. To test for activity after attachment of the sulfhydryl reactive compound, this invention further provides a "small scale" screening technique to readily determine whether modification and attachment has been successful (see Example 9 below).

15 The human IL-3 was additionally modified at its N-terminus in two different and alternative configurations, represented by the "mp" and "m3" designations. The "mp" designation indicates a deletion of the first alanine in the natural human IL-3 protein, thereby changing the N-terminal sequence from MET\*ALA\*PRO to MET\*PRO. The "m3" designation indicates a deletion of the first two amino acids in the natural human IL-3 protein, MET\*ALA\*PRO, to yield a terminus beginning MET\*THR\*GLU\*THR. The reasons  
20 for these modifications have already been discussed. With respect to N-terminus modification of the IL-3 mp $\Delta$ 1Cys19 mutein, amino acids 1-15 were deleted and replaced with the "mp" terminus. The human G-CSF and EPO muteins were additionally modified at the N-terminus by deletion of the first amino acid to obtain "mp" muteins.

It should be understood of course that the depicted list of muteins is merely exemplary and not  
25 exclusive. The design and synthesis of alternative and additional muteins in accord with this invention is well within the present skill in the art. Synthesis of such muteins may be conveniently effected using conventional techniques and methods.

One skilled in the art, of course, could readily design and synthesize other muteins for substitution of cysteine codons or insertion thereof in DNA sequences encoding IL-3, G-CSF and EPO. To modify more  
30 than one site, mutagenesis may be carried out iteratively, or in some cases using an oligonucleotide designed for mutagenesis at more than one site.

#### EXAMPLE 6: Synthesis of DNA molecules encoding CAVs

35 As an alternative to the production of CAV-encoding DNA by mutagenesis of the parental DNA sequence, it should be understood that the desired CAV-encoding DNA may be prepared synthetically. In that case, it will usually be desirable to synthesize the CAV DNA in the form of overlapping oligonucleotides, e.g. overlapping 50-80mers, which together span the desired coding sequence and contain the cysteine additions desired:

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An exemplary EPO mutein made in accordance with this Example is EPO mpCys9. The chemically synthesized cDNA shown in Figure 3 is assembled, modified at position 9, and purified using techniques  
50 know to those skilled in the art. See Wosnick, et al., Gene 60:115-127 (1987); see also U.S. Patent No. 4,904,584. The synthetic cDNA is designed with "overhanging end" nucleotide sequences compatible with those generated by the restriction enzymes, NdeI and XbaI. The purified, synthetically derived, cDNA is ligated with the purified NdeI-XbaI vector portion of plasmid pAL-hIL3-781, resulting in the replacement of human IL-3 cDNA with human erythropoietin cDNA. EPOmpCys166 and EPO mpCys24Cys38Cys83 can be made in the same manner.

55 Given a desired coding sequence, the design, synthesis, assembly and ligation, if desired, to synthetic linkers of other appropriate oligonucleotides is well within the present level of skill in the art.

EXAMPLE 7: PEGylation of the IL-3 mpCys10 mutein

The mutein human IL-3 mpCys10 was prepared in accordance with Example 5 above and PEGylated with two PEG 5000 derivatives, S-Pyridyl Monomethoxy PEG 5000 (PEG 5000 SPDP) and Maleimido Monomethoxy PEG 5000 (PEG 5000 SMCC).

1. PEGylation with S-Pyridyl Monomethoxy PEG 5000: a reducible linkage

## a.) Preparation of the sulfhydryl reactive compound.

PEG 5000 was activated for attachment to a sulfhydryl group as follows. 2.0 grams of Monomethoxy PEG 5000 amine was dissolved in 12 ml dry peroxide free, dioxane. 144 mg (15% excess) of N-succinimidyl-3-(2 pyridyldithio) propionate (SPDP) was added as a dry powder and the reaction was allowed to proceed at room temperature. After 24 hours, the S-pyridyl Monomethoxy PEG 5000 product was precipitated using dry, peroxide free diethyl ether and washed with ether. The product was dried under vacuum to obtain 1.82 grams of white solid, which was identified as S-Pyridyl Monomethoxy PEG 5000 by NMR and IR. The PEG 10,000 analog (PEG 10,000 SPDP) was likewise prepared via an analogous procedure.

## b.) PEGylation of mutein C10 human IL-3.

For this coupling, natural (wild type) human IL-3 was also treated with the PEGylation reagents as a negative control. A stock solution at 1 mg/ml of the mpCys10 mutein in a pH 7 buffered solution of 50 mM  $\text{NaH}_2\text{PO}_4$ , 100 micro M DTT, 1 mM EDTA and about 3mM Guanidine HCl was used. DTT was added to prevent dimerization of the protein; EDTA was added to prevent dimerization via metal mediated oxidative coupling. Guanidine remains as an artifact of the refolding of the protein. A pH 7 was used; a range of 6.5-7.5 is preferred. 0.9 mg of S-Pyridyl Monomethoxy PEG 5000, prepared as set forth above, was weighed into an Eppendorf tube. 360 microliters of the buffered mutein was added and the mixture was vortexed briefly to homogeneity. The reaction was performed at 4 degrees centigrade and when sampled after 2 hours, was found to be complete. Analysis on a 10-20% gradient SDS acrylamide gel stained with Coomassie blue showed the product as nearly pure and running at about 28 kD. (By comparison, mpCys10 and its dimer were used as standards and found to migrate to 15 and 30 kD respectively.) A reducing lane on the gel showed that the PEGylated IL-3 mutein is sensitive to reduction by DTT and regenerated the original protein at about 15 kD.

2. PEGylation with Maleimido Monomethoxy PEG 5000: a non-reducible linkage

## a.) Preparation of the sulfhydryl reactive compound.

In this experiment, PEG 5000 activation was accomplished as follows. 2.0 mg of monomethoxy PEG 5000 amine was dissolved in 12 ml of dry, peroxide free dioxane. 154 mg of sulfosuccinimidyl 4-(N-maleimidomethyl) cyclohexane-1-carboxylate (SMCC) (15% excess) was added as a dry powder and the reaction was allowed to proceed at room temperature. After 24 hours, work up of the product was carried out in the same manner as the S-Pyridyl Monomethoxy PEG 5000 to obtain 1.82 grams Maleimido Monomethoxy PEG 5000. The PEG 10,000 analog was prepared similarly.

## b.) PEGylation of the cys10 IL-3 mutein.

This reaction was carried out in the same manner as the PEGylation reaction using the reducible PEG 5000 reagent and with natural human IL-3 as a negative control. However, 1.0 mg of the PEG derived PEGylating agent Maleimido Monomethoxy PEG 5000 was used. 400 microliters of the mpCys10 IL-3 mutein was added and vortexed to homogeneity. At t=2 hours the reaction was found to be complete. The product was nearly pure and indistinguishable from the S Pyridyl derived conjugate in molecular weight. However, this product is perfectly inert to reductive conditions, such as DTT; in this reducing lane the product, at 28 kD, persists.

In both control reactions, nothing indicative of conjugation is evident at 2 hrs or even at 24 hrs. Selectivity for accessible sulfhydryls in this chemistry is therefore very high.

**EXAMPLE 8: PEGylation of multiple cysteine muteins m3Cys9,10 and m3Cys6,10**

In this experiment, protein stock for both muteins was at 300 µg/ml in the phosphate buffer solution, as described in Example 7. PEGylation stock solutions consisted of the S-pyridyl or Maleimide activated PEG 5000 polymers at 50 µg/ml in the same buffer. To initiate the reaction, 11 µl of the appropriate PEG stock was added to 100 µl of the appropriate protein stock (either the m3Cys9,10 mutein or the m3Cys6,10 mutein) while vortexing. Reactions were allowed to proceed at 4 degrees centigrade overnight. SDS gel analysis of the products as described above revealed that only a trace of starting material remained with both chemistries. Furthermore, both chemistries resulted in new products with a gel mobility of about 37 kD. Reducing lanes on this same gel show that the maleimide conjugate is resistant to reducing, while the S-Pyridyl derived conjugate reverts to starting material.

**EXAMPLE 9: PEGylation of the G-CSF mpAla17Cys37 mutein**

The mutein human G-CSF mpAla17Cys37 was prepared in accordance with Example 5 above and PEGylated with PEG 5000 SPDP, PEG 5000 SMCC and PEG 10,000 SPDP. The natural cysteine at position 17 was deleted and replaced with alanine to prevent possible improper disulfide bridge formation. The mpAla17Cys37 G-CSF protein stock was dialyzed into a pH 7.0 buffering solution of 50mM NaH<sub>2</sub>PO<sub>4</sub> and 1mM EDTA, 100µM DTT and concentrated to approximately 100µg/ml. Total volume was 670 µl.

Stocks of the PEGylation reagents (PEG 5000 SPDP, PEG 5000 SMCC, and PEG 10,000 SPDP) were made up fresh at 10mM in H<sub>2</sub>O (18 x stock solutions prepared as described in Example 7, parts 1a and 1b). The reactions were carried out using the reagents and amounts set forth below.

Rxn#	PEG Reagent	Stock	
		µl Protein	µl PEG
1	5000 SPDP	170	10
2	5000 SMCC	170	10
3	10,000 SPDP	170	10

All reactions were 1mM in PEG reagent. As in Example 7, the mutein was added to the PEG reagent and vortexed to homogeneity. The reactions were allowed to proceed at 4°C overnight and found to be complete.

Analysis on a 10-20% gradient SDS acrylamide gel with both reducing and non-reducing lanes and stained with Coomassie blue showed the mpAla17Cys37 PEG 5000 SPDP and mpAla17Cys37 PEG 5000 SMCC products as nearly pure and running at about 28 kD.

Non-PEGylated G-CSF mutein runs at approximately 19kD. A reducing lane on the gel showed that the SPDP conjugate is sensitive to reduction in the presence of DTT. The SMCC derived reagent was totally resistant to reductive treatment. The PEG 10,000 SPDP PEGylated mpAla17Cys37 G-CSF runs at approximately 32 kD. The non-PEGylated G-CSF mutein is regenerated by DTT treatment.

**EXAMPLE 10: PEGylation of the EPO mpCys9 mutein**

The mutein human EPO mpCys9 prepared in accordance with Example 5 above can be PEGylated using the stock PEGylation reagents prepared in Example 9 and the natural human EPO as a negative control. To initiate the reaction, 10 µl of the appropriate PEG stock as prepared in Example 9 is added to 170 µl of the EPO mpCys9 stock at a concentration of 100µg/ml. The mixture is vortexed and the reaction is allowed to proceed at 4°C overnight. Upon completion, the reaction can be analyzed as in Example 9.

**EXAMPLE 11: Screening of novel CAVs**

Having the constructed novel DNA molecules encoding CAVs in the appropriate expression vector and having attached the sulfhydryl reactive compound to the muteins, it may be desirable to produce each CAV protein on a small scale and "screen" for muteins which possess the desired attachment site or sites. The biological activity of each CAV, before and after attachment of the sulfhydryl reactive compound can be rapidly assessed using an in vitro assay.

Small scale bacterial production of IL-3 CAV muteins

Bacterial strain GI586 was transformed with purified plasmid DNAs consisting of bacterial expression vector, pAL-hIL3-781, ATCC Accession Number 67932, with novel CAV IL-3 coding sequences. The transformed cells were spread on LB agar plates containing 50 µg/ml ampicillin at a density to yield approximately 100 colonies per plate. 3 ml of L broth plus 50 µg/ml ampicillin was inoculated with a single bacterial colony and grown overnight at 30 degrees centigrade. 50 ml of induction media (0.1x L Broth, 1xM9 salts, 0.4% glucose, 1mM MgSO<sub>4</sub>, 50 µg/ml ampicillin) was inoculated with 1 ml of the overnight culture. The 50 ml culture was grown with aeration at 30 degrees centigrade until an 0.5 OD 600 nm level was reached, then the temperature was shifted to 40 degrees centigrade and growth continued for at least 2 hours.

Cells were then harvested by centrifugation at 3500 rpm for 5 minutes in a Sorval centrifuge with a 3B rotor. The supernatant was discarded and the cell pellet resuspended in 1 ml of buffer PED (50 mM NaH<sub>2</sub>PO<sub>4</sub>, pH 7.0, 1 mM EDTA, 5mM DTT, 10 mM PMSF). This 1 ml solution was passed twice through a French Press at 10,000 psi and kept on ice. The solution was then microfuged for 5 minutes at 12,000 rpm. The supernatant was discarded and the pelleted material was resuspended in 150 µl of 7 M guanidine-HCl in PED buffer. The solution was then diluted with 650 µl of PED buffer and placed in dialysis tubing (10,000 MWCO Spectrapore). The sample was dialyzed for at least 4 hours against 2 liters of PED.1 buffer (50 mM NaH<sub>2</sub>PO<sub>4</sub> pH 7.0, 1 mM EDTA, 0.1 mM DTT). The sample was collected and microfuged to remove precipitated proteins. The sample was then analyzed on a 12% Laemmli SDS PAGE gel and the amount of IL-3 protein estimated.

The protein solution was then concentrated to about 0.5 mg/ml, and 200 µg was reacted with a 15 fold molar excess of either S-Pyridyl Monomethoxy PEG 5000 or Maleimido Monomethoxy PEG 5000 for several hours at 4 degrees centigrade. The products were then analyzed by SDS PAGE and biological activity determined by an in vitro TF-1 cell proliferation assay.

This small scale production methodology may be similarly advantageously applied to production of the novel CAV G-CSFs and EPOs of the present invention.

Alternatively, this small scale production for screening may be carried out before attachment of the sulfhydryl reactive compound. In that case, biological activity may still be determined by an in vitro TF-1 cell proliferation assay (or in the case of G-CSF, a 32D cell proliferation assay) and the products may be analyzed by SDS PAGE analysis, in accordance with known techniques.

The same or similar procedures may be used by one skilled in the art to attach other sulfhydryl reactive compounds to the other CAVs of the invention. Homogeneity can be observed by conventional analysis of the modified CAVs so produced e.g. using standard SDS-PAGE or HPLC analysis.

Numerous modifications may be made by one skilled in the art to the methods and compositions of the present invention in view of the disclosure herein. Such modifications are believed to be encompassed by this invention as defined by the appended claims.

**Claims**

1. A cysteine added variant ("CAV") of EPO, characterized in that said CAV comprises a peptide sequence of human EPO modified to contain at least one non-native cysteine residue at which residue said CAV is covalently attached to at least one sulfhydryl reactive compound.
2. A CAV of claim 1, characterized in that the N-terminus of the CAV commences with methionine and the native amino acid at position 1 of the mature peptide sequence is deleted.
3. A CAV of any of claims 1 to 2, characterized in that said at least one sulfhydryl group is selected from one or more of the group consisting of dextran, a carbohydrate based polymer, such as a colominic acid, a polymer of an amino acid, biotin, and a polyalkylene glycol moiety.
4. A CAV of any of claims 1 to 2, characterized in that said at least one sulfhydryl reactive compound is polyethylene glycol.
5. A DNA sequence encoding a CAV comprising a peptide sequence of human EPO modified to contain at least one non-native cysteine residue for attachment of at least one sulfhydryl reactive compound.
6. A host cell containing and capable of expressing a DNA sequence of claim 5.

**EP 0 668 353 A1**

7. A method of producing a CAV of any of claims 1 to 4, comprising covalently attaching at least one  
sulfhydryl reactive compound to at least one non-native cysteine of said CAV produced by culturing a  
host cell containing and capable of expressing a DNA sequence encoding said CAV.
- 5 8. Use of a CAV of any of claims 1 to 4 for the preparation of pharmaceutical compositions suitable for  
stimulating hematopoiesis.

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FIGURE 1

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      1                               10
ATG GCT CCT ATG ACT CAA ACT ACT TCT TTA AAA ACT TCT
Met Ala Pro Met Thr Gln Thr Thr Ser Leu Lys Thr Ser

      25
TGG GTA AAC TGT TCT AAC ATG ATC GAT GAA ATT ATA ACA
Trp Val Asn Cys Ser Asn Met Ile Asp Glu Ile Ile Thr

CAC TTA AAG CAG CCA CCT TTG CCC TTG CTG GAC TTC AAC
His Leu Lys Gln Pro Pro Leu Pro Leu Leu Asp Phe Asn

      40
AAC CTC AAT GGG GAA GAC CAA GAC ATT CTG ATG GAA AAT
Asn Leu Asn Gly Glu Asp Gln Asp Ile Leu Met Glu Asn

      55
AAC CTT CGA AGG CCA AAC CTG GAG GCA TTC AAC AGG GCT
Asn Leu Arg Arg Pro Asn Leu Glu Ala Phe Asn Arg Ala

      70
GTC AAG AGT CTG CAA AAT GCA TCA GCA ATT GAG AGC ATT
Val Lys Ser Leu Gln Asn Ala Ser Ala Ile Glu Ser Ile

      85
CTG AAA AAT CTG CTG CCA TGT CTG CCC CTG GCC ACA GCT
Ile Lys Asn Leu Leu Pro Cys Leu Pro Leu Ala Thr Ala

      100
GCA CCC ACC AGG CAT CCA ATC CAT ATC AAG GAT GGT GAC
Ala Pro Thr Arg His Pro Ile His Ile Lys Asp Gly Asp

      115
TGG AAT GAA TTC CGC CGC AAA CTG ACC TTC TAT CTG AAA
Trp Asn Glu Phe Arg Arg Lys Leu Thr Phe Tyr Leu Lys

ACC CTG GAG AAT GCT CAG GCT CAG CAG ACC ACC CTG AGC
Thr Leu Glu Asn Ala Gln Ala Gln Gln Thr Thr Leu Ser

130
CTC GCG ATC TTC TAG
Leu Ala Ile Phe Stop

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ACC	CCC	CTG	GGC	CCT	GCC	AGC	TCC	CTG	CCC	CAG	AGC	TTC		
Thr	Pro	Leu	Gly	Pro	Ala	Ser	Ser	Leu	Pro	Gln	Ser	Phe		
20														
CTG	CTC	AAG	TGC	TTA	GAG	CAA	GTG	AGG	AAG	ATC	CAG	GGC		
Leu	Leu	Lys	Cys	Leu	Glu	Gln	Val	Arg	Lys	Ile	Gln	Gly		
30														
GAT	GGC	GCA	GCG	CTC	CAG	GAG	AAG	CTG	TGT	GCC	ACC	TAC		
Asp	Gly	Ala	Ala	Leu	Gln	Glu	Lys	Leu	Cys	Ala	Thr	Tyr		
40											50			
AAG	CTG	TGC	CAC	CCC	GAG	GAG	CTG	GTG	CTG	CTC	GGA	CAC		
Lys	Leu	Cys	His	Pro	Glu	Glu	Leu	Val	Leu	Leu	Gly	His		
60														
TCT	CTG	GGC	ATC	CCC	TGG	GCT	CCC	CTG	AGC	AGC	TGC	CCC		
Ser	Leu	Gly	Ile	Pro	Trp	Ala	Pro	Leu	Ser	Ser	Cys	Pro		
70														
AGC	CAG	GCC	CTG	CAG	CTG	GCA	GGC	TGC	TTG	AGC	CAA	CTC		
Ser	Gln	Ala	Leu	Gln	Leu	Ala	Gly	Cys	Leu	Ser	Gln	Leu		
80											90			
CAT	AGC	GGC	CTT	TTC	CTC	TAC	CAG	GGG	CTC	CTG	CAG	GCC		
His	Ser	Gly	Leu	Phe	Leu	Tyr	Gln	Gly	Leu	Leu	Gln	Ala		
100														
CTG	GAA	GGG	ATC	TCC	CCC	GAG	TTG	GGT	CCC	ACC	TTG	GAC		
Leu	Glu	Gly	Ile	Ser	Pro	Glu	Leu	Gly	Pro	Thr	Leu	Asp		
110														
ACA	CTG	CAG	CTG	GAC	GTC	GCC	GAC	TTT	GCC	ACC	ACC	ATC		
Thr	Leu	Gln	Leu	Asp	Val	Ala	Asp	Phe	Ala	Thr	Thr	Ile		
120														
TGG	CAG	CAG	ATG	GAA	GAA	CTG	GGA	ATG	GCC	CCT	GCC	CTG		
Trp	Gln	Gln	Met	Glu	Glu	Leu	Gly	Met	Ala	Pro	Ala	Leu		
140														
CAG	CCC	ACC	CAG	GGT	GCC	ATG	CCG	GCC	TTC	GCC	TCT	GCT		
Gln	Pro	Thr	Gln	Gly	Ala	Met	Pro	Ala	Phe	Ala	Ser	Ala		
150														
TTC	CAG	CGC	CGG	GCA	GGA	GGG	GTC	CTG	GTT	GCC	TCC	CAT		
Phe	Gln	Arg	Arg	Ala	Gly	Gly	Val	Leu	Val	Ala	Ser	His		

FIGURE 2b

				160									
CTG	CAG	AGC	TTC	CTG	GAG	GTG	TCG	TAC	CGC	GTT	CTA	CGC	
Leu	Gln	Ser	Phe	Leu	Glu	Val	Ser	Tyr	Arg	Val	Leu	Arg	
170				174									
CAC	CTT	GCC	CAG	CCC	T								
His	Leu	Ala	Gln	Pro									

21

FIGURE 3b

Lys	Glu	Ala	Ile	Ser	Pro	Pro	Asp	Ala	Ala	Ser	Ala	Ala
AAG	GAA	GCC	ATC	TCC	CCT	CCA	GAT	GCG	GCC	TCA	GCT	GCT
TTC	CTT	CGG	TAG	AGG	GGA	GGT	CTA	CGC	CGG	AGT	CGA	CGA

130

Pro	Leu	Arg	Thr	Ile	Thr	Ala	Asp	Thr	Phe	Arg	Lys	Leu
CCA	CTC	CGA	ACA	ATC	ACT	GCT	GAC	ACT	TTC	CGA	AAA	CTC
GGT	GAG	GCT	TGT	TAG	TGA	CGA	CTG	TGA	AAG	GCT	TTT	GAG

145

Phe	Arg	Val	Tyr	Ser	Asn	Phe	Leu	Arg	Gly	Lys	Leu	Lys
TTC	CGA	GTC	TAC	TCC	AAT	TTC	CTC	CGG	GGA	AAG	CTG	AAG
AAG	GCT	CAG	ATG	AGG	TTA	AAG	GAG	GCC	CCT	TTC	GAC	TTC

160

Leu	Tyr	Thr	Gly	Glu	Ala	Cys	Arg	Thr	Gly	Asp	Arg
CTG	TAC	ACA	GGG	GAG	GCC	TGC	AGG	ACA	GGG	GAC	AGA
GAC	ATG	TGT	CCC	CTC	CGG	ACG	TCC	TGT	CCC	CTG	TCT

STOP  
TAA TAATGATAGGATCCT  
ATT ATTACTATCCTAGGAGATC - 5'



European Patent  
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## EUROPEAN SEARCH REPORT

Application Number  
EP 95 10 3989

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P,Y	EP-A-0 357 804 (GENETICS INSTITUTE INC. CETUS CORPORATION) * the whole document *	1-8	
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The present search report has been drawn up for all claims			
Place of search		Date of completion of the search	Examiner
THE HAGUE		21 June 1995	Chambonnet, F
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X : particularly relevant if taken alone Y : particularly relevant if combined with another document of the same category A : technological background O : non-written disclosure P : intermediate document		T : theory or principle underlying the invention E : earlier patent document, but published on, or after the filing date D : document cited in the application L : document cited for other reasons A : member of the same patent family, corresponding document	

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